#6/a

SEQUENCE LISTING

<110> Detmar, Michael J.
Streit, Michael

<120> THROMBOSPONDIN-2 AND USES THEREOF

<130> 10287-051001

<140> 09/536,087
<141> 2000-03-24

<150> 60/127,221
<151> 1999-03-31

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ctg tgg gtg tgg ccc agc acg caa gct ggt cac cag gac aaa gac acg      100
Leu Trp Val Trp Pro Ser Thr Gln Ala Gly His Gln Asp Lys Asp Thr
 10                  15                  20                  25

acc ttc gac ctt ttc agt atc agc aac atc aac cgc aag acc att ggc      148
Thr Phe Asp Leu Phe Ser Ile Ser Asn Ile Asn Arg Lys Thr Ile Gly
                 30                  35                  40

gcc aag cag ttc cgc ggg ccc gac ccc ggc gtg ccg gct tac cgc ttc      196
Ala Lys Gln Phe Arg Gly Pro Asp Pro Gly Val Pro Ala Tyr Arg Phe
             45                  50                  55

gtg cgc ttt gac tac atc cca ccg gtg aac gca gat gac ctc agc aag      244
Val Arg Phe Asp Tyr Ile Pro Pro Val Asn Ala Asp Asp Leu Ser Lys
         60                  65                  70

atc acc aag atc atg cgg cag aag gag ggc ttc ttc ctc acg gcc cag      292
Ile Thr Lys Ile Met Arg Gln Lys Glu Gly Phe Phe Leu Thr Ala Gln
     75                  80                  85

ctc aag cag gac ggc aag tcc agg ggc acg ctg ttg gct ctg gag ggc      340
Leu Lys Gln Asp Gly Lys Ser Arg Gly Thr Leu Leu Ala Leu Glu Gly
 90                  95                  100                 105
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ccc ggt ctc tcc cag agg cag ttc gag atc gtc tcc aat ggc ccc gcg      388
Pro Gly Leu Ser Gln Arg Gln Phe Glu Ile Val Ser Asn Gly Pro Ala
                110                 115                 120

gac acg ctg gat ctc acc tac tgg att gac ggc acc cgg cat gtg gtc      436
Asp Thr Leu Asp Leu Thr Tyr Trp Ile Asp Gly Thr Arg His Val Val
            125                 130                 135

tcc ctg gag gac gtc ggc ctg gct gac tcg cag tgg aag aac gtc acc      484
Ser Leu Glu Asp Val Gly Leu Ala Asp Ser Gln Trp Lys Asn Val Thr
        140                 145                 150

gtg cag gtg gct ggc gag acc tac agc ttg cac gtg ggc tgc gac ctc      532
Val Gln Val Ala Gly Glu Thr Tyr Ser Leu His Val Gly Cys Asp Leu
    155                 160                 165

ata gac agc ttc gct ctg gac gag ccc ttc tac gag cac ctg cag gcg      580
Ile Asp Ser Phe Ala Leu Asp Glu Pro Phe Tyr Glu His Leu Gln Ala
170                 175                 180                 185

gaa aag agc cgg atg tac gtg gcc aaa ggc tct gcc aga gag agt cac      628
Glu Lys Ser Arg Met Tyr Val Ala Lys Gly Ser Ala Arg Glu Ser His
                190                 195                 200

ttc agg ggt ttg ctt cag aac gtc cac cta gtg ttt gaa aac tct gtg      676
Phe Arg Gly Leu Leu Gln Asn Val His Leu Val Phe Glu Asn Ser Val
            205                 210                 215

gaa gat att cta agc aag aag ggt tgc cag caa ggc cag gga gct gag      724
Glu Asp Ile Leu Ser Lys Lys Gly Cys Gln Gln Gly Gln Gly Ala Glu
        220                 225                 230

atc aac gcc atc agt gag aac aca gag acg ctg cgc ctg ggt ccg cat      772
Ile Asn Ala Ile Ser Glu Asn Thr Glu Thr Leu Arg Leu Gly Pro His
    235                 240                 245

gtc acc acc gag tac gtg ggc ccc agc tca gag agg agg ccc gag gtg      820
Val Thr Thr Glu Tyr Val Gly Pro Ser Ser Glu Arg Arg Pro Glu Val
250                 255                 260                 265

tgc gaa cgc tcg tgc gag gag ctg gga aac atg gtc cag gag ctc tcg      868
Cys Glu Arg Ser Cys Glu Glu Leu Gly Asn Met Val Gln Glu Leu Ser
                270                 275                 280

ggg ctc cac gtc ctc gtg aac cag ccc agc gag aac ctc aag aga gtg      916
Gly Leu His Val Leu Val Asn Gln Pro Ser Glu Asn Leu Lys Arg Val
            285                 290                 295

tcg aat gat aac cag ttt ctc tgg gag ctc att ggt ggc cct cct aag      964
Ser Asn Asp Asn Gln Phe Leu Trp Glu Leu Ile Gly Gly Pro Pro Lys
        300                 305                 310

aca agg aac atg tca gct tgc tgg cag gat ggc cgg ttc ttt gcg gaa     1012
Thr Arg Asn Met Ser Ala Cys Trp Gln Asp Gly Arg Phe Phe Ala Glu
    315                 320                 325
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aat gaa acg tgg gtg gtg gac agc tgc acc acg tgt acc tgc aag aaa     1060
Asn Glu Thr Trp Val Val Asp Ser Cys Thr Thr Cys Thr Cys Lys Lys
330                 335                 340                 345

ttt aaa acc att tgc cac caa atc acc tgc ccg cct gca acc tgc gcc     1108
Phe Lys Thr Ile Cys His Gln Ile Thr Cys Pro Pro Ala Thr Cys Ala
                350                 355                 360

agt cca tcc ttt gtg gaa ggc gaa tgc tgc cct tcc tgc ctc cac tcg     1156
Ser Pro Ser Phe Val Glu Gly Glu Cys Cys Pro Ser Cys Leu His Ser
            365                 370                 375

gtg gac ggt gag gag ggc tgg tct ccg tgg gca gag tgg acc cag tgc     1204
Val Asp Gly Glu Glu Gly Trp Ser Pro Trp Ala Glu Trp Thr Gln Cys
        380                 385                 390

tcc gtg acg tgt ggc tct ggg acc cag cag aga ggc cgg tcc tgt gac     1252
Ser Val Thr Cys Gly Ser Gly Thr Gln Gln Arg Gly Arg Ser Cys Asp
    395                 400                 405

gtc acc agc aac acc tgc ttg ggg ccc tcc atc cag aca cgg gct tgc     1300
Val Thr Ser Asn Thr Cys Leu Gly Pro Ser Ile Gln Thr Arg Ala Cys
410                 415                 420                 425

agt ctg agc aag tgt gac acc cgc atc cgg cag gac ggc ggc tgg agc     1348
Ser Leu Ser Lys Cys Asp Thr Arg Ile Arg Gln Asp Gly Gly Trp Ser
                430                 435                 440

cac tgg tca cct tgg tct tca tgc tct gtg acc tgt gga gtt ggc aat     1396
His Trp Ser Pro Trp Ser Ser Cys Ser Val Thr Cys Gly Val Gly Asn
            445                 450                 455

atc aca cgc atc cgt ctc tgc aac tcc cca gtg ccc cag atg ggg ggc     1444
Ile Thr Arg Ile Arg Leu Cys Asn Ser Pro Val Pro Gln Met Gly Gly
        460                 465                 470

aag aat tgc aaa ggg agt ggc cgg gag acc aaa gcc tgc cag ggc gcc     1492
Lys Asn Cys Lys Gly Ser Gly Arg Glu Thr Lys Ala Cys Gln Gly Ala
    475                 480                 485

cca tgc cca atc gat ggc cgc tgg agc ccc tgg tcc ccg tgg tcg gcc     1540
Pro Cys Pro Ile Asp Gly Arg Trp Ser Pro Trp Ser Pro Trp Ser Ala
490                 495                 500                 505

tgc act gtc acc tgt gcc ggt ggg atc cgg gag cgc acc cgg gtc tgc     1588
Cys Thr Val Thr Cys Ala Gly Gly Ile Arg Glu Arg Thr Arg Val Cys
                510                 515                 520

aac agc cct gag cct cag tac gga ggg aag gcc tgc gtg ggg gat gtg     1636
Asn Ser Pro Glu Pro Gln Tyr Gly Gly Lys Ala Cys Val Gly Asp Val
            525                 530                 535

cag gag cgt cag atg tgc aac aag agg agc tgc ccc gtg gat ggc tgt     1684
Gln Glu Arg Gln Met Cys Asn Lys Arg Ser Cys Pro Val Asp Gly Cys
        540                 545                 550

tta tcc aac ccc tgc ttc ccg gga gcc cag tgc agc agc ttc ccc gat     1732
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Leu Ser Asn Pro Cys Phe Pro Gly Ala Gln Cys Ser Ser Phe Pro Asp
    555                 560                 565

ggg tcc tgg tca tgc ggc tcc tgc cct gtg ggc ttc ttg ggc aat ggc     1780
Gly Ser Trp Ser Cys Gly Ser Cys Pro Val Gly Phe Leu Gly Asn Gly
570                 575                 580                 585

acc cac tgt gag gac ctg gac gag tgt gcc ctg gtc ccc gac atc tgc     1828
Thr His Cys Glu Asp Leu Asp Glu Cys Ala Leu Val Pro Asp Ile Cys
                590                 595                 600

ttc tcc acc agc aag gtg cct cgc tgt gtc aac act cag cct ggc ttc     1876
Phe Ser Thr Ser Lys Val Pro Arg Cys Val Asn Thr Gln Pro Gly Phe
            605                 610                 615

cac tgc ctg ccc tgc ccg ccc cga tac aga ggg aac cag ccc gtc ggg     1924
His Cys Leu Pro Cys Pro Pro Arg Tyr Arg Gly Asn Gln Pro Val Gly
        620                 625                 630

gtc ggc ctg gaa gca gcc aag acg gaa aag caa gtg tgt gag ccc gaa     1972
Val Gly Leu Glu Ala Ala Lys Thr Glu Lys Gln Val Cys Glu Pro Glu
    635                 640                 645

aac cca tgc aag gac aag aca cac aac tgc cac aag cac gcg gag tgc     2020
Asn Pro Cys Lys Asp Lys Thr His Asn Cys His Lys His Ala Glu Cys
650                 655                 660                 665

atc tac ctg ggc cac ttc agc gac ccc atg tac aag tgc gag tgc cag     2068
Ile Tyr Leu Gly His Phe Ser Asp Pro Met Tyr Lys Cys Glu Cys Gln
                670                 675                 680

aca ggc tac gcg ggc gac ggg ctc atc tgc ggg gag gac tcg gac ctg     2116
Thr Gly Tyr Ala Gly Asp Gly Leu Ile Cys Gly Glu Asp Ser Asp Leu
            685                 690                 695

gac ggc tgg ccc aac ctc aat ctg gtc tgc gcc acc aac gcc acc tac     2164
Asp Gly Trp Pro Asn Leu Asn Leu Val Cys Ala Thr Asn Ala Thr Tyr
        700                 705                 710

cac tgc atc aag gat aac tgc ccc cat ctg cca aat tct ggg cag gaa     2212
His Cys Ile Lys Asp Asn Cys Pro His Leu Pro Asn Ser Gly Gln Glu
    715                 720                 725

gac ttt gac aag gac ggg att ggc gat gcc tgt gat gat gac gat gac     2260
Asp Phe Asp Lys Asp Gly Ile Gly Asp Ala Cys Asp Asp Asp Asp Asp
730                 735                 740                 745

aat gac ggt gtg acc gat gag aag gac aac tgc cag ctc ctc ttc aat     2308
Asn Asp Gly Val Thr Asp Glu Lys Asp Asn Cys Gln Leu Leu Phe Asn
                750                 755                 760

ccc cgc cag gct gac tat gac aag gat gag gtt ggg gac cgc tgt gac     2356
Pro Arg Gln Ala Asp Tyr Asp Lys Asp Glu Val Gly Asp Arg Cys Asp
            765                 770                 775

aac tgc cct tac gtg cac aac cct gcc cag atc gac aca gac aac aat     2404
Asn Cys Pro Tyr Val His Asn Pro Ala Gln Ile Asp Thr Asp Asn Asn
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        780                 785                 790

gga gag ggt gac gcc tgc tcc gtg gac att gat ggg gac gat gtc ttc     2452
Gly Glu Gly Asp Ala Cys Ser Val Asp Ile Asp Gly Asp Asp Val Phe
    795                 800                 805

aat gaa cga gac aat tgt ccc tac gtc tac aac act gac cag agg gac     2500
Asn Glu Arg Asp Asn Cys Pro Tyr Val Tyr Asn Thr Asp Gln Arg Asp
810                 815                 820                 825

acg gat ggt gac ggt gtg ggg gat cac tgt gac aac tgc ccc ctg gtg     2548
Thr Asp Gly Asp Gly Val Gly Asp His Cys Asp Asn Cys Pro Leu Val
                830                 835                 840

cac aac cct gac cag acc gac gtg gac aat gac ctt gtt ggg gac cag     2596
His Asn Pro Asp Gln Thr Asp Val Asp Asn Asp Leu Val Gly Asp Gln
            845                 850                 855

tgt gac aac aac gag gac ata gat gac gac ggc cac cag aac aac cag     2644
Cys Asp Asn Asn Glu Asp Ile Asp Asp Asp Gly His Gln Asn Asn Gln
        860                 865                 870

gac aac tgc ccc tac atc tcc aac gcc aac cag gct gac cat gac aga     2692
Asp Asn Cys Pro Tyr Ile Ser Asn Ala Asn Gln Ala Asp His Asp Arg
    875                 880                 885

gac ggc cag ggc gac gcc tgt gac cct gat gat gac aac gat ggc gtc     2740
Asp Gly Gln Gly Asp Ala Cys Asp Pro Asp Asp Asp Asn Asp Gly Val
890                 895                 900                 905

ccc gat gac agg gac aac tgc cgg ctt gtg ttc aac cca gac cag gag     2788
Pro Asp Asp Arg Asp Asn Cys Arg Leu Val Phe Asn Pro Asp Gln Glu
                910                 915                 920

gac ttg gac ggt gat gga cgg ggt gat att tgt aaa gat gat ttt gac     2836
Asp Leu Asp Gly Asp Gly Arg Gly Asp Ile Cys Lys Asp Asp Phe Asp
            925                 930                 935

aat gac aac atc cca gat att gat gat gtg tgt cct gaa aac aat gcc     2884
Asn Asp Asn Ile Pro Asp Ile Asp Asp Val Cys Pro Glu Asn Asn Ala
        940                 945                 950

atc agt gag aca gac ttc agg aac ttc cag atg gtc ccc ttg gat ccc     2932
Ile Ser Glu Thr Asp Phe Arg Asn Phe Gln Met Val Pro Leu Asp Pro
    955                 960                 965

aaa ggg acc acc caa att gat ccc aac tgg gtc att cgc cat caa ggc     2980
Lys Gly Thr Thr Gln Ile Asp Pro Asn Trp Val Ile Arg His Gln Gly
970                 975                 980                 985

aag gag ctg gtt cag aca gcc aac tcg gac ccc ggc atc gct gta ggt     3028
Lys Glu Leu Val Gln Thr Ala Asn Ser Asp Pro Gly Ile Ala Val Gly
                990                 995                1000

ttt gac gag ttt ggg tct gtg gac ttc agt ggc aca ttc tac gta aac     3076
Phe Asp Glu Phe Gly Ser Val Asp Phe Ser Gly Thr Phe Tyr Val Asn
            1005                1010                1015
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act gac cgg gac gac gac tat gcc ggc ttc gtc ttt ggt tac cag tca     3124
Thr Asp Arg Asp Asp Asp Tyr Ala Gly Phe Val Phe Gly Tyr Gln Ser
        1020                1025                1030

agc agc cgc ttc tat gtg gtg atg tgg aag cag gtg acg cag acc tac     3172
Ser Ser Arg Phe Tyr Val Val Met Trp Lys Gln Val Thr Gln Thr Tyr
    1035                1040                1045

tgg gag gac cag ccc acg cgg gcc tat ggc tac tcc ggc gtg tcc ctc     3220
Trp Glu Asp Gln Pro Thr Arg Ala Tyr Gly Tyr Ser Gly Val Ser Leu
1050                1055                1060                1065

aag gtg gtg aac tcc acc acg ggg acg ggc gag cac ctg agg aac gcg     3268
Lys Val Val Asn Ser Thr Thr Gly Thr Gly Glu His Leu Arg Asn Ala
                1070                1075                1080

ctg tgg cac acg ggg aac acg ccg ggg cag gtg cga acc tta tgg cac     3316
Leu Trp His Thr Gly Asn Thr Pro Gly Gln Val Arg Thr Leu Trp His
            1085                1090                1095

gac ccc agg aac att ggc tgg aag gac tac acg gcc tat agg tgg cac     3364
Asp Pro Arg Asn Ile Gly Trp Lys Asp Tyr Thr Ala Tyr Arg Trp His
        1100                1105                1110

ctg act cac agg ccc aag acc ggc tac atc aga gtc tta gtg cat gaa     3412
Leu Thr His Arg Pro Lys Thr Gly Tyr Ile Arg Val Leu Val His Glu
    1115                1120                1125

gga aaa cag gtc atg gca gac tca gga cct atc tat gac caa acc tac     3460
Gly Lys Gln Val Met Ala Asp Ser Gly Pro Ile Tyr Asp Gln Thr Tyr
1130                1135                1140                1145

gct ggc ggg cgg ctg ggt cta ttt gtc ttc tct caa gaa atg gtc tat     3508
Ala Gly Gly Arg Leu Gly Leu Phe Val Phe Ser Gln Glu Met Val Tyr
                1150                1155                1160

ttc tca gac ctc aag tac gaa tgc aga gat att taaacaagat ttgctgcatt   3561
Phe Ser Asp Leu Lys Tyr Glu Cys Arg Asp Ile
            1165                1170

tccggcaatg ccctgtgcat gccatggtcc ctaga                              3596

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<212> PRT
<213> Homo sapiens

<400> 2
Met Val Trp Arg Leu Val Leu Leu Ala Leu Trp Val Trp Pro Ser Thr
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Gln Ala Gly His Gln Asp Lys Asp Thr Thr Phe Asp Leu Phe Ser Ile
            20                  25                  30
Ser Asn Ile Asn Arg Lys Thr Ile Gly Ala Lys Gln Phe Arg Gly Pro
        35                  40                  45
Asp Pro Gly Val Pro Ala Tyr Arg Phe Val Arg Phe Asp Tyr Ile Pro
    50                  55                  60
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Pro Val Asn Ala Asp Asp Leu Ser Lys Ile Thr Lys Ile Met Arg Gln
65                  70                  75                  80
Lys Glu Gly Phe Phe Leu Thr Ala Gln Leu Lys Gln Asp Gly Lys Ser
                85                  90                  95
Arg Gly Thr Leu Leu Ala Leu Glu Gly Pro Gly Leu Ser Gln Arg Gln
            100                 105                 110
Phe Glu Ile Val Ser Asn Gly Pro Ala Asp Thr Leu Asp Leu Thr Tyr
        115                 120                 125
Trp Ile Asp Gly Thr Arg His Val Val Ser Leu Glu Asp Val Gly Leu
    130                 135                 140
Ala Asp Ser Gln Trp Lys Asn Val Thr Val Gln Val Ala Gly Glu Thr
145                 150                 155                 160
Tyr Ser Leu His Val Gly Cys Asp Leu Ile Asp Ser Phe Ala Leu Asp
                165                 170                 175
Glu Pro Phe Tyr Glu His Leu Gln Ala Glu Lys Ser Arg Met Tyr Val
            180                 185                 190
Ala Lys Gly Ser Ala Arg Glu Ser His Phe Arg Gly Leu Leu Gln Asn
        195                 200                 205
Val His Leu Val Phe Glu Asn Ser Val Glu Asp Ile Leu Ser Lys Lys
    210                 215                 220
Gly Cys Gln Gln Gly Gln Gly Ala Glu Ile Asn Ala Ile Ser Glu Asn
225                 230                 235                 240
Thr Glu Thr Leu Arg Leu Gly Pro His Val Thr Thr Glu Tyr Val Gly
                245                 250                 255
Pro Ser Ser Glu Arg Arg Pro Glu Val Cys Glu Arg Ser Cys Glu Glu
            260                 265                 270
Leu Gly Asn Met Val Gln Glu Leu Ser Gly Leu His Val Leu Val Asn
        275                 280                 285
Gln Pro Ser Glu Asn Leu Lys Arg Val Ser Asn Asp Asn Gln Phe Leu
    290                 295                 300
Trp Glu Leu Ile Gly Gly Pro Pro Lys Thr Arg Asn Met Ser Ala Cys
305                 310                 315                 320
Trp Gln Asp Gly Arg Phe Phe Ala Glu Asn Glu Thr Trp Val Val Asp
                325                 330                 335
Ser Cys Thr Thr Cys Thr Cys Lys Lys Phe Lys Thr Ile Cys His Gln
            340                 345                 350
Ile Thr Cys Pro Pro Ala Thr Cys Ala Ser Pro Ser Phe Val Glu Gly
        355                 360                 365
Glu Cys Cys Pro Ser Cys Leu His Ser Val Asp Gly Glu Glu Gly Trp
    370                 375                 380
Ser Pro Trp Ala Glu Trp Thr Gln Cys Ser Val Thr Cys Gly Ser Gly
385                 390                 395                 400
Thr Gln Gln Arg Gly Arg Ser Cys Asp Val Thr Ser Asn Thr Cys Leu
                405                 410                 415
Gly Pro Ser Ile Gln Thr Arg Ala Cys Ser Leu Ser Lys Cys Asp Thr
            420                 425                 430
Arg Ile Arg Gln Asp Gly Gly Trp Ser His Trp Ser Pro Trp Ser Ser
        435                 440                 445
Cys Ser Val Thr Cys Gly Val Gly Asn Ile Thr Arg Ile Arg Leu Cys
    450                 455                 460
Asn Ser Pro Val Pro Gln Met Gly Gly Lys Asn Cys Lys Gly Ser Gly
465                 470                 475                 480
Arg Glu Thr Lys Ala Cys Gln Gly Ala Pro Cys Pro Ile Asp Gly Arg
                485                 490                 495
Trp Ser Pro Trp Ser Pro Trp Ser Ala Cys Thr Val Thr Cys Ala Gly
            500                 505                 510
Gly Ile Arg Glu Arg Thr Arg Val Cys Asn Ser Pro Glu Pro Gln Tyr
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        515                 520                 525
Gly Gly Lys Ala Cys Val Gly Asp Val Gln Glu Arg Gln Met Cys Asn
    530                 535                 540
Lys Arg Ser Cys Pro Val Asp Gly Cys Leu Ser Asn Pro Cys Phe Pro
545                 550                 555                 560
Gly Ala Gln Cys Ser Ser Phe Pro Asp Gly Ser Trp Ser Cys Gly Ser
                565                 570                 575
Cys Pro Val Gly Phe Leu Gly Asn Gly Thr His Cys Glu Asp Leu Asp
            580                 585                 590
Glu Cys Ala Leu Val Pro Asp Ile Cys Phe Ser Thr Ser Lys Val Pro
        595                 600                 605
Arg Cys Val Asn Thr Gln Pro Gly Phe His Cys Leu Pro Cys Pro Pro
    610                 615                 620
Arg Tyr Arg Gly Asn Gln Pro Val Gly Val Gly Leu Glu Ala Ala Lys
625                 630                 635                 640
Thr Glu Lys Gln Val Cys Glu Pro Glu Asn Pro Cys Lys Asp Lys Thr
                645                 650                 655
His Asn Cys His Lys His Ala Glu Cys Ile Tyr Leu Gly His Phe Ser
            660                 665                 670
Asp Pro Met Tyr Lys Cys Glu Cys Gln Thr Gly Tyr Ala Gly Asp Gly
        675                 680                 685
Leu Ile Cys Gly Glu Asp Ser Asp Leu Asp Gly Trp Pro Asn Leu Asn
    690                 695                 700
Leu Val Cys Ala Thr Asn Ala Thr Tyr His Cys Ile Lys Asp Asn Cys
705                 710                 715                 720
Pro His Leu Pro Asn Ser Gly Gln Glu Asp Phe Asp Lys Asp Gly Ile
                725                 730                 735
Gly Asp Ala Cys Asp Asp Asp Asp Asp Asn Asp Gly Val Thr Asp Glu
            740                 745                 750
Lys Asp Asn Cys Gln Leu Leu Phe Asn Pro Arg Gln Ala Asp Tyr Asp
        755                 760                 765
Lys Asp Glu Val Gly Asp Arg Cys Asp Asn Cys Pro Tyr Val His Asn
    770                 775                 780
Pro Ala Gln Ile Asp Thr Asp Asn Asn Gly Glu Gly Asp Ala Cys Ser
785                 790                 795                 800
Val Asp Ile Asp Gly Asp Asp Val Phe Asn Glu Arg Asp Asn Cys Pro
                805                 810                 815
Tyr Val Tyr Asn Thr Asp Gln Arg Asp Thr Asp Gly Asp Gly Val Gly
            820                 825                 830
Asp His Cys Asp Asn Cys Pro Leu Val His Asn Pro Asp Gln Thr Asp
        835                 840                 845
Val Asp Asn Asp Leu Val Gly Asp Gln Cys Asp Asn Asn Glu Asp Ile
    850                 855                 860
Asp Asp Asp Gly His Gln Asn Asn Gln Asp Asn Cys Pro Tyr Ile Ser
865                 870                 875                 880
Asn Ala Asn Gln Ala Asp His Asp Arg Asp Gly Gln Gly Asp Ala Cys
                885                 890                 895
Asp Pro Asp Asp Asp Asn Asp Gly Val Pro Asp Asp Arg Asp Asn Cys
            900                 905                 910
Arg Leu Val Phe Asn Pro Asp Gln Glu Asp Leu Asp Gly Asp Gly Arg
        915                 920                 925
Gly Asp Ile Cys Lys Asp Asp Phe Asp Asn Asp Asn Ile Pro Asp Ile
    930                 935                 940
Asp Asp Val Cys Pro Glu Asn Asn Ala Ile Ser Glu Thr Asp Phe Arg
945                 950                 955                 960
Asn Phe Gln Met Val Pro Leu Asp Pro Lys Gly Thr Thr Gln Ile Asp
                965                 970                 975
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Pro Asn Trp Val Ile Arg His Gln Gly Lys Glu Leu Val Gln Thr Ala
            980                 985                 990
Asn Ser Asp Pro Gly Ile Ala Val Gly Phe Asp Glu Phe Gly Ser Val
        995                 1000                1005
Asp Phe Ser Gly Thr Phe Tyr Val Asn Thr Asp Arg Asp Asp Asp Tyr
    1010                1015                1020
Ala Gly Phe Val Phe Gly Tyr Gln Ser Ser Ser Arg Phe Tyr Val Val
1025                1030                1035                1040
Met Trp Lys Gln Val Thr Gln Thr Tyr Trp Glu Asp Gln Pro Thr Arg
                1045                1050                1055
Ala Tyr Gly Tyr Ser Gly Val Ser Leu Lys Val Val Asn Ser Thr Thr
            1060                1065                1070
Gly Thr Gly Glu His Leu Arg Asn Ala Leu Trp His Thr Gly Asn Thr
        1075                1080                1085
Pro Gly Gln Val Arg Thr Leu Trp His Asp Pro Arg Asn Ile Gly Trp
    1090                1095                1100
Lys Asp Tyr Thr Ala Tyr Arg Trp His Leu Thr His Arg Pro Lys Thr
1105                1110                1115                1120
Gly Tyr Ile Arg Val Leu Val His Glu Gly Lys Gln Val Met Ala Asp
                1125                1130                1135
Ser Gly Pro Ile Tyr Asp Gln Thr Tyr Ala Gly Gly Arg Leu Gly Leu
            1140                1145                1150
Phe Val Phe Ser Gln Glu Met Val Tyr Phe Ser Asp Leu Lys Tyr Glu
        1155                1160                1165
Cys Arg Asp Ile
    1170

<210> 3
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<212> PRT
<213> Homo sapiens

<400> 3
Asp Lys Asp Thr Thr Phe Asp Leu Phe Ser Ile Ser Asn Ile Asn
 1               5                  10                  15

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<211> 27
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<220>
<223> primer for PCR

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gaattcagga gctcagctgc aggaggc                                          27

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<220>
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gaattctagg gaccatggca tgcac                                            25
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<212> PRT
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<220>
<223> Synthetically generated peptide

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Arg Glu Ser His Phe Arg Gly Leu Leu Gln Asn Val His Leu Val Phe
 1   5   10   15

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<211> 18
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetically generated peptide

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 1   5   10   15
Ser Cys

<210> 8
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<220>
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 1   5   10   15
Cys Lys Lys Phe Lys Thr
            20

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<212> PRT
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<220>
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Glu Leu Ile Gly Gly Pro Pro Lys Thr Arg Asn Met Ser Ala Cys
 1   5   10   15

<210> 10
<211> 7
<212> PRT
<213> Artificial Sequence

<220>
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<400> 10
Trp Ser Pro Trp Ala Glu Trp
 1   5

<210> 11
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<212> PRT
<213> Homo sapiens

<400> 11
Cys Ser Val Thr Val Gly
 1   5